OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/841,321A

DATE: 07/09/2001 TIME: 13:25:23

Input Set: A:\BERL 020-04 corrected.txt
Output Set: N:\CRF3\07092001\1841321A.raw

3 <110> APPLICANT: Urry, Dan 5 <120> TITLE OF INVENTION: Injectable Implants For Tissue Augmentation and Restoration 7 <130> FILE REFERENCE: BERL-020/04US 9 <140> CURRENT APPLICATION NUMBER: 09/841,321A C--> 10 <141> CURRENT FILING DATE: 2001-04-30 12 <150> PRIOR APPLICATION NUMBER: US 09/258,723 13 <151> PRIOR FILING DATE: 1999-02-26 15 <150> PRIOR APPLICATION NUMBER: US 60/087155 16 <151> PRIOR FILING DATE: 1998-05-29 18 <150> PRIOR APPLICATION NUMBER: US 60/076297 19 <151> PRIOR FILING DATE: 1998-02-27 ENTERED 21 <160> NUMBER OF SEQ ID NOS: 65 23 <170> SOFTWARE: PatentIn version 3.0 25 <210> SEO ID NO: 1 26 <211> LENGTH: 180 27 <212> TYPE: DNA 28 <213> ORGANISM: Artificial Sequence 30 <220> FEATURE: 31 <221> NAME/KEY: misc feature 32 <222> LOCATION: (1)..(180) 33 <223> OTHER INFORMATION: Synthetic 36 <400> SEQUENCE: 1 37 gaggateega agacaacagg tggtgtteeg ggeggegtae egggtggegt aeegggeggt 60 39 ttcccgggag gtgtgccggg tggggttcca ggcggtgtac cgggtgggtt tccqggcggt 120 41 gttccgggtg gagttccggg tggcgtgccg ggcggttttc caggaagtct tcggatccag 180 44 <210> SEQ ID NO: 2 45 <211> LENGTH: 113 46 <212> TYPE: DNA 47 <213> ORGANISM: Artificial Sequence 49 <220> FEATURE: 50 <221> NAME/KEY: misc feature 51 <222> LOCATION: (1)..(113) 52 <223> OTHER INFORMATION: Synthetic 55 <400> SEQUENCE: 2 56 gaggatecag gegttggggt acegggtgtt ggegtaeegg gtaaaggtgt eeegggegtt 58 ggtgtgccgg gtgtaggctt tccgggtttc ggattcccag gcgttggatc cag 113 61 <210> SEQ ID NO: 3 62 <211> LENGTH: 33 63 <212> TYPE: DNA 64 <213> ORGANISM: Artificial Sequence 66 <220> FEATURE: 67 <221> NAME/KEY: misc\_feature 68 <222> LOCATION: (1)..(33) 69 <223> OTHER INFORMATION: Synthetic 72 <400> SEQUENCE: 3 73 taggggtacc gggtcgtggt gactctccgg gcg 33

76 <210> SEQ ID NO: 4

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Input Set : A:\BERL 020-04 corrected.txt Output Set: N:\CRF3\07092001\1841321A.raw

77 (011) I DNOW! 22	
77 <211> LENGTH: 33	
78 <212> TYPE: DNA	
79 <213> ORGANISM: Artificial Sequence 81 <220> FEATURE:	
82 <221> NAME/KEY: misc_feature	
83 <222> LOCATION: (1)(33)	
84 <223> OTHER INFORMATION: Synthetic	
87 <400> SEQUENCE: 4	
88 cgcatcccca tggcccagca ccactgagag gcc	33
91 <210> SEQ ID NO: 5	
92 <211> LENGTH: 111	
93 <212> TYPE: DNA	
94 <213> ORGANISM: Artificial Sequence	
96 <220> FEATURE:	
97 <221> NAME/KEY: misc_feature	
98 <222> LOCATION: (1)(111)	
99 <223> OTHER INFORMATION: Synthetic	
102 <400> SEQUENCE: 5	
103 gaggatccag gcgttggggt accgggtgtt ggcgtaccgg gtgttggtgt cccgggcaaa	60
105 ggtgtgccgg gtgtaggcgt tccgggtgtg ggagtcccag gcgttggatc c	111
108 <210> SEQ ID NO: 6	
109 <211> LENGTH: 345	
110 <212> TYPE: DNA	
111 <213> ORGANISM: Artificial Sequence	
113 <220> FEATURE:	
114 <221> NAME/KEY: misc_feature	
115 <222> LOCATION: (1)(345)	
116 <223> OTHER INFORMATION: Synthetic	
119 <400> SEQUENCE: 6	
120 ggcgttggtg taccgggtgt tggtgtgccg ggtgttggtg ttccgggcgt aggcgtaccg	60
122 ggcgtaggcg tgccgggcgt aggcgttccg ggcgtgggcg taccgggcgt gggcgtgccg	120
124 ggtgtgggcg tecegggtgt aggtgtteca ggcgtagggg tacegggteg tggtgaetet	180
126 ccgggcgttg gtgtaccggg tgttggtgtg ccgggtgttg gtgttccggg cgtaggcgta	240
128 ccgggcgtag gcgtgccggg cgtaggcgtt ccgggcgtgg gcgtaccggg cgtgggcgtg	300
130 ccgggtgtgg gcgtcccggg tgtaggtgtt ccaggcgttg gatcc	345
133 <210> SEQ ID NO: 7 134 <211> LENGTH: 463	•
·	
135 <212> TYPE: DNA	
136 <213> ORGANISM: Artificial Sequence 138 <220> FEATURE:	
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140 <222> LOCATION: (1)(463)	
140 <222> LOCATION: (1)(463) 141 <223> OTHER INFORMATION: Synthetic	
144 <400> SEQUENCE: 7	
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145 ggatccaggc gttggtgtac cgggtgttgg tgtgccgggt gttggtgttc cgggcgtagg	60
147 cgtaccgggc gtaggcgtgc cgggcgtagg cgttccgggc gtaggcgtac cgggcgtggg	120
149 cgtgccgggt gtgggcgtcc caggtgtagg cgttccgggt gtgggtgtag ctccgggtgt	180 240
151 taggegttage cegggegtag gtgttgetee gggegttage gtggegeegg gtgttggtgt	300
153 tgctccgggt gtaggcgttg ctccgggcgt tggtgttgcc ccaggtgtag gtgtggcacc	200

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155 gggcgttggt gtaccgggtg ttggtgtgcc gggtgttggt gttccgggcg taggcgtacc
                                                                           360
157 gggcgtaggc gtgccgggcg taggcgttcc gggcgtgggc gtaccgggcg tgggcgtgcc
                                                                           420
159 gggtgtgggc gtcccgggtg taggtgttcc aggcgttgga tcc
                                                                           463
162 <210> SEQ ID NO: 8
163 <211> LENGTH: 111
164 <212> TYPE: DNA
165 <213> ORGANISM: Artificial Sequence
167 <220> FEATURE:
168 <221> NAME/KEY: misc feature
169 <222> LOCATION: (1)..(111)
170 <223> OTHER INFORMATION: Synthetic
173 <400> SEQUENCE: 8
174 gaggatccag gcgttggggt accgggtgtt ggcgtaccgg gtgaaggtgt cccgggcgtt
                                                                            60
176 ggtgtgccgg gtgtaggcgt tccgggtgtg ggagtcccag gcgttggatc c
                                                                           111
179 <210> SEQ ID NO: 9
180 <211> LENGTH: 48
181 <212> TYPE: PRT
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <221> NAME/KEY: PEPTIDE
186 <222> LOCATION: (1)..(48)
187 <223> OTHER INFORMATION: Synthetic
190 <400> SEQUENCE: 9
192 Gly Gly Val Pro Gly Gly Val Pro Gly Gly Val Pro Gly Gly Phe Pro
193 1
                    5
                                         10
195 Gly Gly Val Pro Gly Gly Val Pro Gly Gly Val Pro Gly Gly Phe Pro
                20
                                     25
198 Gly Gly Val Pro Gly Gly Val Pro Gly Gly Val Pro Gly Gly Phe Pro
199
            35
                                 40
201 <210> SEQ ID NO: 10
202 <211> LENGTH: 30
203 <212> TYPE: PRT
204 <213> ORGANISM: Artificial Sequence
206 <220> FEATURE:
207 <221> NAME/KEY: PEPTIDE
208 <222> LOCATION: (1)..(30)
209 <223> OTHER INFORMATION: Synthetic
212 <400> SEQUENCE: 10
214 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Lys Gly Val Pro Gly
217 Val Gly Val Pro Gly Val Gly Phe Pro Gly Phe Gly Phe Pro
218
                20
                                     25
220 <210> SEQ ID NO: 11
221 <211> LENGTH: 11
222 <212> TYPE: PRT
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <221> NAME/KEY: PEPTIDE
227 <222> LOCATION: (1)..(11)
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228 <223> OTHER INFORMATION: Synthetic
231 <400> SEQUENCE: 11
233 Val Gly Val Pro Gly Arg Gly Asp Ser Pro Gly
234 1
236 <210> SEQ ID NO: 12
237 <211> LENGTH: 30
238 <212> TYPE: PRT
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <221> NAME/KEY: PEPTIDE
243 <222> LOCATION: (1)..(30)
244 <223> OTHER INFORMATION: Synthetic
247 <400> SEQUENCE: 12
249 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Lys Gly Val Pro Gly
250 1
252 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
                                    25
253
                20
255 <210> SEQ ID NO: 13
256 <211> LENGTH: 111
257 <212> TYPE: PRT
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <221> NAME/KEY: PEPTIDE
262 <222> LOCATION: (1)..(111)
263 <223> OTHER INFORMATION: Synthetic
266 <400> SEOUENCE: 13
268 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
271 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
274 Gly Val Pro Gly Val Gly Val Pro Gly Val Pro Gly Val Gly
277 Val Pro Gly Val Gly Val Pro Gly Arg Gly Asp Ser Pro Gly Val Gly
                            55
280 Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
283 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
                    8.5
286 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
                100
289 <210> SEQ ID NO: 14
290 <211> LENGTH: 148
291 <212> TYPE: PRT
292 <213> ORGANISM: Artificial Sequence
294 <220> FEATURE:
295 <221> NAME/KEY: PEPTIDE
296 <222> LOCATION: (1)..(148)
297 <223> OTHER INFORMATION: Synthetic
300 <400> SEQUENCE: 14
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```
302 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
305 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val
                                    25
308 Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly
311 Val Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val
314 Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro
315 65
                        70
                                            75
317 Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly Val
320 Ala Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val
321
                100
                                    105
323 Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
324
                                120
            115
326 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro Gly
327 130
                            135
329 Val Gly Val Pro
330 145
332 <210> SEQ ID NO: 15
333 <211> LENGTH: 30
334 <212> TYPE: PRT
335 <213> ORGANISM: Artificial Sequence
337 <220> FEATURE:
338 <221> NAME/KEY: PEPTIDE
339 <222> LOCATION: (1)..(30)
340 <223> OTHER INFORMATION: Synthetic
343 <400> SEQUENCE: 15
345 Gly Val Gly Val Pro Gly Val Gly Val Pro Gly Glu Gly Val Pro Gly
348 Val Gly Val Pro Gly Val Gly Val Pro Gly Val Gly Val Pro
349
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351 <210> SEQ ID NO: 16
352 <211> LENGTH: 4
353 <212> TYPE: PRT
354 <213> ORGANISM: Artificial Sequence
356 <220> FEATURE:
357 <221> NAME/KEY: PEPTIDE
358 <222> LOCATION: (1)..(4)
359 <223> OTHER INFORMATION: Synthetic
362 <400> SEQUENCE: 16
364 Val Pro Gly Gly
365 1
367 <210> SEQ ID NO: 17
368 <211> LENGTH: 5
369 <212> TYPE: PRT
370 <213> ORGANISM: Artificial Sequence
372 <220> FEATURE:
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Use of n and/or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

DATE: 07/09/2001

PATENT APPLICATION: US/09/841,321A

TIME: 13:25:24

Input Set : A:\BERL 020-04 corrected.txt
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 $L:10 \ M:271 \ C:$  Current Filing Date differs, Replaced Current Filing Date

L:657 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19